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Scoring table: Sequence: Tabular output Run on: MPsrch\_pp \* Release 3.1A John F. Collins, Biocomputing Research Unit. Copyright (c) 1993-1998 University of Edinburgh, U.K. Distribution rights by Oxford Molecular Ltd protein - protein database search, using Smith-Waterman algorithm not generated. >US-09-331-631-1 (186-248) from US09331631.pep (5 of 5) 494 1 KRDPQQREYEDCRRRCEQQE.....MMNPQRGGSGRYEEGEEEQS 63 Sat May 13 09:51:03 2000; MasPar time 4.55 Seconds 327.787 Million cell updates/sec (MT)

Description: Perfect Score: Title:

PAM 150 Gap 11

188963 seqs, 23686106 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Database: a-geneseq35 1:geneseqp

Statistics: Mean 25.007; Variance 101.250; scale 0.247

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

22		21	20	19	18	17	16	15	14	13	12	11	10	9	œ	7	თ	υı	4	ω	2	سو	Result No. So
	80	81	81	81	84	85	87	88	92	93	99	110	110	110	115	141	157	170	170	433	449	494	Score
	16.2	16.4	16.4	16.4		17.2	17.6	17.8	18.6	18.8	20.0	22.3	22.3	22.3	23.3	28.5	31.8	34.4	34.4	87.7	90.9	100.0	Query Match
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W62836		W14987	W31867	W03474	W33628	W73369	W30750	W30749	W37483	W62841	P90996	P93109	W14783	R12223	W62835	W62837	W62832	R20181	W62831	W62830	W62829	W62828	ID
	Zea mays antimicrobial	Protein derived from r	Human metastasis-assoc	Mouse SRY-related prot	Yeast transcriptional	77	Partial human YT521 ge	Rat YT521 gene product	Mouse liver cancer-ori			Human androgen recepto	Androgen receptor.	Human androgen recepto	Zea mays antimicrobial	Hordeum vulgare antimi	Gossypium hirsutum ant	encoded	Theobroma cacao antimi	Macadamia integrifolia	Macadamia integrifolia	Macadamia integrifolia	Description
	.06e+01	1.71e+01	1.71e+01	1.71e+01	9.75e+00		5:52e+00		2.11e+00					•			3.22e-06		1.94e-07	.40e-	.23e-	7.86e-40	Pred. No.

RESULT
ID W6
AC W6
DT 27
DE Ma

w62829 standard; Protein; 666 AA.
w62829;
27-OCT-1998 (first entry)
Macadamia integrifolia antimicrobial protein.
antimiorobial protein; infestation; control.

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186 KRDPQQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEE 245

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45	44	43	42	41	40	39	38	37	36	35	34	<b>ω</b>	32	31	30	29	28	27	26	25	24
76	76	76	77	77	77	77	77	77	78	78	78	78	79	79	79	79	08	08	80	80	08
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W78191	Y04889	Y04890	W52822	P61362	P71081	W26536	W52825	W06970	W26407	W92415	R49790	W08136	W48896	R89214	R41875	W52828	R71111	R82630	W48796	W03662	W95073
Human secreted protein	Mycobacterium species	Mycobacterium species	Human NonO/TFE3 fusion	Soybean glycinin A3B4	Sequence encoded by Vi	Trypanosoma cruzi anti	Human TFE3/NonO 5'-RAC	Modified hsp60 gene pr	Matrix/scaffold-associ	Human SATB1 protein.	Sequence of special AT	Human cytokine respons	Candida albicans CaCLA	Peroxisome proliferato	Steroid hormone recept	Human NonO DNA protein	Spinocerebellar ataxia	70K autoantigen, part	Drosphila melanogaster	Human 70K U1 snRNP pro	GST-HD fusion protein
4.32e+01	4.32e+01	4.32e+01	3.59e+01	3.59e+01	3.59e+01	3.59e+01	3.59e+01	3.59e+01	2.99e+01	2.99e+01	2.99e+01	2.99e+01	2.48e+01	2.48e+01	2.48e+01	2.48e+01	2.06e+01	2.06e+01	2.06e+01	2.06e+01	2.06e+01

## ALIGNMENTS

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Query Match Best Local Similarity Matches 63; Conse	be used to control microbial infestations animals. Sequence 666 AA;	Claim 1; Page 34-36; 95pp; English. The sequence is that of an antimicrobial protein which can	useful for controlling microbial infestations of	N-PSDB; V42310.	WPI; 98-377279/32.	Bower NI, Goulter	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY		٠.	02-JUL-1998.	WO9827805-A1.		Protein		Peptide	Key	Macadamia integ	antimicrobial protein;	Macadamia integrifolia antimicrobial	27-OCT-1998 (f	••	JLT 1 W62828 standard;	
100.0%; larity 100.0%; Conservative	AA; .	s that of an	trolling micr		32.	ter KC, Green	ES CENT TROPI	AU-004275.	AU0874.			/note= "mature	29666	<pre>/note= "signal peptide"</pre>	128	Location/Qualifiers	integrifolia.	protein; infe	grifolia anți	(first entry)		d; Protein; 666	
Score 49 Pred. No 0; Mis	l infesta	English. antimicro	obial inf	5 6 1 1		JL, Manı	CAL PLANT					ure protein"		nal pepti		alifiers		infestation;	microbial			66 AA.	
Score 494; DB 1; Pred. No. 7.86e-40; 0; Mismatches 0	ations in	obial prot	microbial infestations of	1		Green JL, Manners JM, Marcus	r PATHOLOG					ein"		ide"				control.	l protein.				
	plant	ein v	of			larcus																	
Length 666;	in plants and mammalian	which can	plants or ma			з JP;																	
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Query Match
Best Local S
Matches 5
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Best Local :
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                                                                                                                                                         Novel anti-microbial protein from e.g. Macadamia integruseful for controlling microbial infestations of plants Claim 1; Page 43-45; 96pp; English.

The sequence is that of an antimicrobial protein which be used to control microbial infestations in plants and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man Claim 1; Page 39-41; 96pp; English.

The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammal be used to control microbial infestations in plants.
                                                                                                                                                                                                                                      02-JUL-1998.
22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, Goulter KC, Green JL, Manners JM, Mar
                                                                                                                                                                                                                                                                                                                                                                                                                       Macadamia integrifolia antimicrobial protein. antimicrobial protein; infestation; control.
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22-DEC-1997; AU0874.
22-DEC-1996; AU-004275.
(RETR.) COOP RES CENT TROPICAL |
BOWET NI, GOULTER KC, Green JL,
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W62830 standard; Protein; 625
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les 58; Conser
                     KRDPQQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGSGRYEEGEE 204
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larity 88.9%;
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                                                                           Score 433; DB 1;
Pred. No. 1.40e-33;
3; Mismatches 4
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Pred. No. 3.23e-35;
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Cocoa; fl
                                                                                      Claim 4; Fig 2; 59pp; English.

The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocoa beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologies between the 67 kD polypeptide and the vicilins, which are seed storage proteins.
                                                                                                                                                                                                                                                                         LI-JUN-1990; GB-013016.

(MRSC ) MARS UK LTD.

Spencer ME, Hodge R, Deakin
WPI; 92-024418/03.

N-PSDB- 020777
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20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPIC
Bower NI, Goulter KC, Green
WPI; 98-377279/32.
                                                                                                                                                                                                                           beans and produced expression vectors
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R20181;
16-APR-1992 (fi
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The sequence is that of an antimicrobial protein which.
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beans and produced in large c
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11-JUN-1990; GB-013(
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            be used to control microbial infestations in plants and mammalian animals.
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QRQYQQCQGRCQEQQQGQREQQQCQRKCWEQYKEQERGEHENYHNHK-KNRSEEEEEGQQ 139
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re encoded by 67 kD T. cacao protein flavour; vicilin; seed storage protein
                                   Similarity 40.78;
24; Conservative
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1. No. 1.94e-07;
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    02-JUL-1998.
22-DEC-1997;
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(RETR-) COOP
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22-DEC-1997;
20-DEC-1996;
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
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(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
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WPI; 98-377279/32.

Novel anti-microbial protein from e.g. Macadamia integruseful for controlling microbial infestations of plants Claim 1; Page 49-51; 96pp; English.

The sequence is that of an antimicrobial protein which be used to control microbial infestations in plants and
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22-DEC-1997;
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Zea mays.
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(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
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23; Conservative
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RES CENT TROPICAL PLANT PATHOLOGY.
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Pred. No. 9.77e-05;
12; Mismatches 18
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d. No. 3.22e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence was deduced from a cDNA clone isolated by screening commercially available human testis and prostate lambda gtll cDNA libraries. The sequence is very similar to that of rat AR and in the DNA-binding domain it is identical to that of rAR DNA-binding domain. Homology comparisons with other known steroid receptors indicate that hAR is more closely related to glucocorticoid, mineralo-corticoid and progesterone receptors than to v-erb-A or treceptors for oestrogen, vitamin D and thyroid hormones.
                                                                                     Homo sapiens.
W09711170-A1
                                                                                                                           Androgen receptor.
Androgen receptor; acidic fibroblast growth factor; aFGF; antisense: beniqn prostatic hyperplasia; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30~MAY~1991.
19-OCT~1990; 1
17-NOV~1989;
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The sequence is that of an antimicrobial protein
27-MAR-1997. \( \)
20-SEP-1996; U15081. \( \)
20-SEP-1995; US-004018. \( \)
(WORC-) WORCESTER FOUND
                                                                                                                                                                                                                   22-JUN-1997
                                                                                                                                                                                                                                                  W14783;
                                                                                                                                                                                                                                                                         W14783
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This sequence was deduced from a cDNA clone isolated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ARCH-) ARCH DEV CORP.
Liao S, Chang C;
WPI; 91-178048/24.
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R12223;
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                                                                                                                                                                                                                                                                                                                                                                        EQQEPRQOHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ
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                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 22.3%;
Similarity 39.1%;
18; Conservative
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US-438775.
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/label= DNA-binding do
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                                                                                                                                                                                                                                                                      Protein;
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Pred. No. 2.17e-02;
19; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 110; DB 1; Le
Pred. No. 5.99e-02;
11; Mismatches 16;
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Human androgen receptor (W14/83) binds testosterone and, acting at the transcriptional level, regulates the growth of normal prostatic cells. Antisense oligonucleotides (see also T63200, T63404-05) based on an androgen receptor cDNA clone (see also T63407) can be used to prevent androgen receptor gene expression, the treatment of benign prostatic hyperplasia and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA encoding androgen receptor protein - useful for transforming eukaryotic hosts for protein expression and subsequent antibody prodn. Disclosure; Fig. 4; 41p; English.

Androgen receptor protein (AR) is used to produce mono- or poly-clonal antibodies. These are used for the detection and quantification of AR in the presence of endogenous androgen, as androgen will not interfere with binding. They may be used in assays to determine and quantify cellular distribution of AR in tumour tissue, and are esp. useful for evaluating prostate cancers to determine responsiveness to androgen withdrawal
                                                                                                                                                                                                                                                              JT 12
P90996 standard; protein;
P90996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-APR-1989; U01548.
14-APR-1988; US-182646.
(UYNC-) University of North Carolina.
French FS, Wilson EM, Joseph DR, Lubahn DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zamecnik PA;
WPI; 97-202879/18.
N-PSDB; T63407.
     region
                                                                                                                    Key
                                                                                                                                                                      Human androgen receptor DNA clone. Androgen receptor; TR2 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 89-324206/44.
N-PSDB; N91772.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens. WO8909791-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human androgen receptor; ployclonal antibody; cancer.
                                                                                       region
                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                28-FEB-1990 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human androgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P93109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P93109
                                                                                                                                                                                                                                                                                                                                                                                                    202 EQQEPROQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         202 EQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                              8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83
                                                                                                                                                                                                                                                                                                                                                                                                                                                         OQQQQQQQQETSPRQQQQQQGEDGSPQAHRRGPTG-YLVLDEEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QQQQQQQQQQETSPRQQQQQQGEDGSPQAHRRGPTG-YLVLDEEQ 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; protein; 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 22.3%;
Similarity 39.1%;
18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
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     /*tag= a
/product=98 kD polypeptide
185. .919
                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22.3%;
                                                                                                                                                                                                                                                                                        919
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Pred. No. 5.99e-02;
11; Mismatches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 110; DB 1; Le
Pred. No. 5.99e-02;
11; Mismatches 16;
                                                                                                                                                                                                                                                                                        AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
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RESULT

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Best Local S
Matches 1
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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W09827805-A1.

02-JUL-1998.

22-DEC-1997; AU0874.

20-DEC-1996; AU-004275.
                                      09-DEC-1997.
27-MAY-1996;
27-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                W37483 stand
W37483;
27-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO8909223-A.
05-OCT-1989.
24-MAR-1989;
30-MAR-1988;
                                                                                                                                                            Mus sp.
J09313185-A.
                                                                                                                                                                                                                                             Mouse; human;
hHDGF; HET-A;
                                                                                                                                                                                                                                                                                   Mouse liver cancer-originated culture Mouse; human; liver cancer-originated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 66; 96pp; Eng
The sequence is that of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RETR-), COOP RES CENT TROPICAL PLANT PATHOLOGY BOWER NI, GOULTER KC, Green JL, Manners JM, May WPI; 98-377279/32 WPI; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stenocarpus sinuatus antimicrobial protein. antimicrobial protein; infestation; control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W62841;
27-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The protein is used to raise antibodies for receptor assays and for affinity purification.
The 98 kD product starts at the first Met codon; the 79 kD product starts from the second.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             purification.
Claim 8; fig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New DNA encoding new androgen receptor and TR2 polypeptide(s) - to bind DNA, and derived antibodies, useful for receptor assay a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ARCH-) Arch Development
Liao S, Chang C;
WPI; 89-309501/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and man
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W62841 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB
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|: |: ||:|| || ||:||
REYEDCRRRCEQQE--PRQQHQC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; N91577
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    KISHIMOTO
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Similarity 39.1%;
18; Conservative
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                                          131788.
JP-131788.
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US-176107.
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10; M
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1. No. 5.38e-01;
Mismatches 17;
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1.74e+00;
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PT polypeptide of at least 5 continuous amino acids, useful in producing, e.g. protein HET-A

PT producing, e.g. protein HET-A

PS Claim 10; Page 13-14; 18pp; Japanese.

CC The present sequence represents a protein which has local homology to a human liver cancer-originated culture cell isolated from a mouse testicle cDNA library. The sequence may (1) be modified to produce a collypeptide comprising at least 5 continuous amino acids; (2) a collypeptide comprising at the sequence of a collypeptide protein similar to protein of (1), but comprises no protein combined naturally and also has 2 amino acid sequences (as given in the specification) which may be modified to produce a sequence of at cleast 5 amino acids; (3) a recombinant DNA molecule comprising a vector and the DNA segment of (1); (4) a cell comprising (3); (5) producing a plypeptide comprising a protein of (1), to produce the polypeptide which is isolated. The above method may be used to produce proteins HET-A and HET-B.
                                                                                                                                        Query Match 17.8%;
Best Local Similarity 28.6%;
Matches 16; Conservative
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Best Local Similarity 35.6%;
Matches 21; Conservative
                                                                                                                                                                                                                                                                            Brain ischaemia-related gene and protein - useful for elucidation of mechanisms of ischaemia and in animal models for screening drugs for prevention or treatment of ischaemia
Claim 1; Pages 10-12; 14pp; Japanese.
The present sequence is the rat YT521 gene product. The YT521 gene is expressed specifically upon re-oxygenation of astrocytes after exposure to low-oxygen conditions. The protein is involved in brain ischaemia, so it and the related gene are useful for elucidation of the mechanism of ischaemia. They are also useful for the creation of a model animal with diseases accompanying ischaemia or for development of drugs for prevention or treatment of such diseases. Antibodies to the protein are useful for determining the severity of diseases caused by ischaemia. Sequence 712 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J09238685-A.
16-SEP-1997.
07-MAR-1996; 049380.
07-MAR-1996; JP-049380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat YT521 gene product.
rat; YT521 gene; re-oxygenation; astrocyte; low-oxygen condition; brain; ischaemia; animal model; drug screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W30749 standard; Protein; 712 AA W30749;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TANA ) TANABE SEIYAKU CO.
(TOYA/) TOYAMA M.
WPI; 97-506554/47.
                                  N-PSDB; V01731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              322 RRD-EERRRELEARRRREQEE-ELRRLREQEREE-KERRKERAERGGSSG-EELEDEEP 376
186 KRD-PQQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 92; DB 1; Len
Pred. No. 2.11e+00;
17; Mismatches 16;
                                                                                                                                        Score 88; DB 1; Le
Pred. No. 4.56e+00;
19; Mismatches 16;
                                                                                                                                                                                                        Length 712;
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